CHICKEN c-SRC cDNA

(SEQ ID NO:2)

1 tctgacaccc atctgtctgt ctgtctgtgt gctgcaggag ctgagctgac tctgctgtgg
61 cctcgcgtac cactgtggcc aggcggtagc tgggacgtgc agcccaccac catggggagc
121 agcaagagca agcccaagga ccccagccag cgccggcgca gcctggagcc acccgacagc
181 acccaccacg ggggattccc agcctcgcag acccccaaca agacagcagc ccccgacacg
241 caccgcaccc ccagccgctc ctttgggacc gtggccaccg agcccaagct cttcgggggc
301 ttcaacactt ctgacaccgt tacgtcgccg cagcgtgccg gggcactggc tggcggcgtc
361 accactttcg tggctctcta cgactacgag tcccggactg aaacggactt gtccttcaag
421 aaaggagaac gcctgcagat tgtcaacaac acggaaggtg actggtggct ggctcattcc
481 ctcactacag gacagacggg ctacatcccc agtaactatg tcgcgccctc agactccatc
541 caggctgaag agtggtactt tgggaagatc actcgtcggg agtccgagcg gctgctgctc
601 aaccccgaaa accccccggg aaccttcttg gtccgggaga gcgagacgac aaaaggtgcc
661 tattgcctct ccgtttctga ctttgacaac gccaaggggc tcaatgtgaa gcactacaag
721 atccgcaagc tggacagcgg cggcttctac atcacctcac gcacacagtt cagcagcctg
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841 gtctgcccca cgtccaagcc ccagacccag ggactcgcca aggacgcgtg ggaaatcccc
901 cgggagtcgc tgcggctgga ggtgaagctg gggcagggct gctttggaga ggtctggatg
961 gggacctgga acggcaccac cagagtggcc ataaagactc tgaagcccgg caccatgtcc
1021 ccggaggcct tcctgcagga agcccaagtg atgaagaagc tccggcatga gaagctggtt
1081 cagctgtacg cagtggtgtc ggaagagccc atctacatcg tcactgagta catgagcaag
1141 gggagcctcc tggatttcct gaagggagag atgggcaagt acctgcggct gccacagctc
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1441 gatgtctggt ccttcggcat cctgctgact gagctgacca ccaagggccg ggtgccatac
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1561 ccgcccgagt gccccgagtc gctgcatgac ctcatgtgcc agtgctggcg gagggaccct
1621 gaggagcggc ccacttttga gtacctgcag gccttcctgg aggactactt cacctcgaca
1681 gagccccagt accagcctgg agagaaccta taggcctgga gctcctcctg gaccagaggc
1741 ctcgctgtgg ggtacaggg

FIG. 1

CHICKEN cSRC ENCODED PROTEIN

(SEQ ID NO:3)

MGSSKSKPKDPSQRRRSLEPPDSTHHGGFPASQTPNKTAA

PDTHRTPSRSFGTVATEPKLFGGFNTSDTVTSPQRAGALA

GGVTTFVALYDYESRTETDLSFKKGERLQIVNNTEGDWWL

AHSLTTGQTGYIPSNYVAPSDSIQAEEWYFGKITRRESER

LLLNPENPRGTFLVRESETTKGAYCLSVSDFDNAKGLNVK

HYKIRKLDSGGFYITSRTQFSSLQQLVAYYSKHADGLCHR

LTNVCPTSKPQTQGLAKDAWEIPRESLRLEVKLGQGCFGE

VWMGTWNGTTRVAIKTLKPGTMSPEAFLQEAQVMKKLRHE

KLVQLYAVVSEEPIYIVTEYMSKGSLLDFLKGEMGKYLRL

PQLVDMAAQIASGMAYVERMNYVHRDLRAANILVGENL

VCKVADFGLARLIEDNEYTARQGAKFPIKWTAPEAALYGR

FTIKSDVWSFGILLTELTTKGRVPYPGMVNREVLDQVERG

YRMPCPPECPESLHDLMCQCWRRDPEERPTFEYLQAFLE

DYFTSTEPQYQPGENL

FIG. 2

HUMAN c-SRC cDNA

(SEQ ID NO:4)

1 gcgccgcgtc ccgcaggccg tgatgccgcc cgcgcggagg tggccccgac cgcagtgccc
61 caagagagct ctaatggtac caagtgacag gttggcttta ctgtgactcg gggacgccag
121 agctcctgag aagatgtcag caatacaggc cgcctggcca tccggtacag aatgtattgc
181 caagtacaac ttccacggca ctgccgagca ggacctgccc ttctgcaaag gagacgtgct
241 caccattgtg gccgtcacca aggaccccaa ctggtacaaa gccaaaaaca aggtgggccg
301 tgagggcatc atcccagcca actacgtcca gaagcgggag ggcgtgaagg cgggtaccaa
361 actcagcctc atgccttggt tccacggcaa gatcacacgg gagcaggctg agcggcttct
421 gtacccgccg gagacaggcc tgttcctggt gcgggagagc accaactacc ccggagacta
481 cacgctgtgc gtgagctgcg acggcaaggt ggagcactac cgcatcatgt accatgccag
541 caagctcagc atcgacgagg aggtgtactt tgagaacctc atgcagctgg tggagcacta
601 cacctcagac gcagatggac tctgtacgcg cctcattaaa ccaaaggtca tggagggcac
661 agtggcggcc caggatgagt tctaccgcag cggctgggcc ctgaacatga aggagctgaa
721 gctgctgcag accatcggga agggggagtt cggagacgtg atgctgggcg attaccgagg
781 gaacaaagtc gccgtcaagt gcattaagaa cgacgccact gcccaggcct tcctggctga
841 agcctcagtc atgacgcaac tgcggcatag caacctggtg cagctcctgg gcgtgatcgt
901 ggaggagaag ggcgggctct acatcgtcac tgagtacatg gccaagggga gccttgtgga
961 ctacctgcgg tctaggggtc ggtcagtgct gggcggagac tgtctcctca agttctcgct
1021 agatgtctgc gaggccatgg aatacctgga gggcaacaat ttcgtgcatc gagacctggc
1081 tgcccgcaat gtgctggtgt ctgaggacaa cgtggccaag gtcagcgact ttggtctcac
1141 caaggaggcg tccagcaccc aggacacggg caagctgcca gtcaagtgga cagcccctga
1201 ggccctgaga gagaagaaat tctccactaa gtctgacgtg tggagtttcg gaatccttct
1261 ctgggaaatc tactcctttg ggcgagtgcc ttatccaaga attccccctga aggacgtcgt
1321 ccctcgggtg gagaagggct acaagatgga tgccccccgac ggctgccccgc ccgcagtcta
1381 tgaagtcatg aagaactgct ggcacctgga cgccgccatg cggccctcct tcctacagct
1441 ccgagagcag cttgagcaca tcaaaaccca cgagctgcac ctgtgacggc tggcctccgc
1501 ctgggtcatg ggcctgtggg gactgaacct ggaagatcat ggacctggtg ccccctgctca
1561 ctgggcccga gcctgaactg agccccagcg ggctggcggg ccttttcct gcgtcccagc
1621 ctgcaccct ccggcccgt ctctcttgga cccacctgtg gggcctgggg agcccactga
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1741 ctggcctccc gccactcgcc ttcttagagt tttattcctt tccttttttg agatttttt
1801 tccgtgtgtt tatttttat tatttttcaa gataaggaga aagaaagtac ccagcaaatg
1861 ggcattttac aagaagtacg aatcttattt ttcctgtcct gcccgtgagg gtgggggga
1921 ccgggccccct ctctagggac ccctcgcccc agcctcattc cccattctgt gtcccatgtc
1981 ccgtgtctcc tcggtcgccc cgtgtttgcg cttgaccatg ttgcactgtt tgcatgcgcc
2041 cgaggcagac gtctgtcagg ggcttggatt tcgtgtgccg ctgccacccg cccacccgcc
2101 ttgtgagatg gaattgtaat aaaccacgcc atgaggacac cgccgcccgc ctcggcgctt
2161 cctccaccga aaaaaaaaaa aaaaaaa

FIG. 3

HUMAN c-SRC ENCODED PROTEIN

(SEQ ID NO:5)

MSAIQAAWPSGTECIAKYNFHGTAEQDLPFCKGDVLTIVAVTKD

PNWYKAKNKVGREGIIPANYVQKREGVKAGTKLSLMPWFHGKIT

REQAERLLYPPETGLFLVRESTNYPGDYTLCVSCDGKVEHYRIMY

HASKLSIDEEVYFENLMQLVEHYTSDADGLCTRLIKPKVMEGTVA

AQDEFYRSGWALNMKELKLLQTIGKGEFGDVMLGDYRGNKVAV

KCIKNDATAQAFLAEASVMTQLRHSNLVQLLGVIVEEKGGLYIVTE

YMAKGSLVDYLRSRGRSVLGGDCLLKFSLDVCEAMEYLEGNNFVH

RDLAARNVLVSEDNVAKVSDFGLTKEASSTQDTGKLPVKWTAPEAL

REKKFSTKSDVWSFGILLWEIYSFGRVPYPRIPLKDVVPRVEKGYKM

DAPDGCPPAVYEVMKNCWHLDAAMRPSFLQLREQLEHIKTHELHL

FIG. 4

Activation of endogenous Src activity by bFGF and VEGF

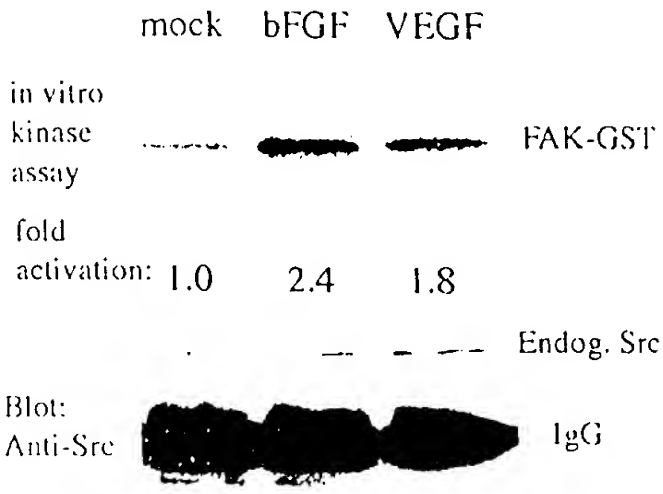
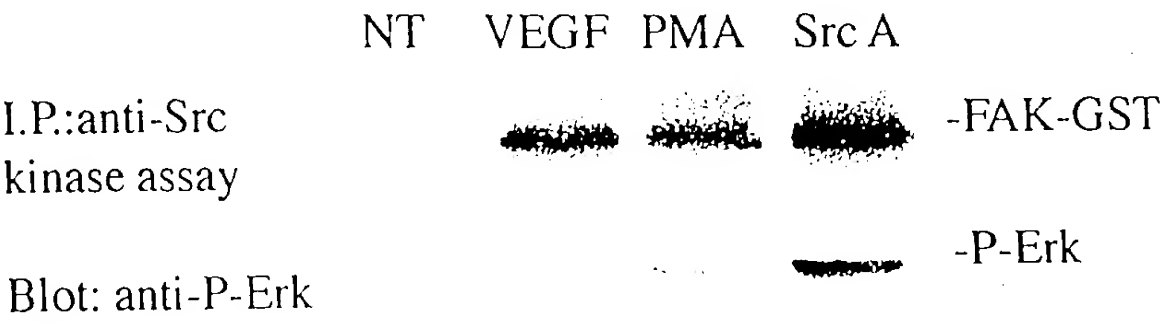


FIG. 5

Retroviral expression of Src A activates vascular MAP kinase phosphorylation



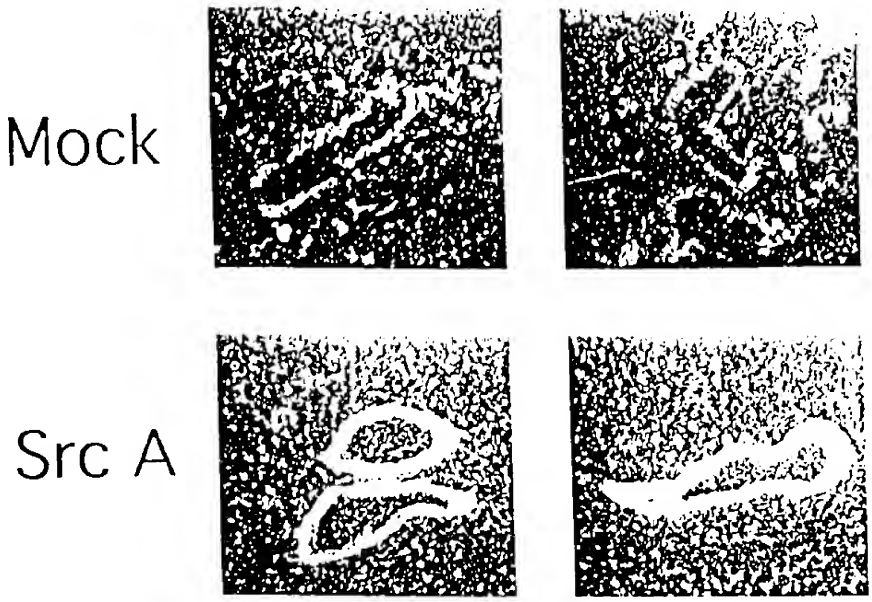
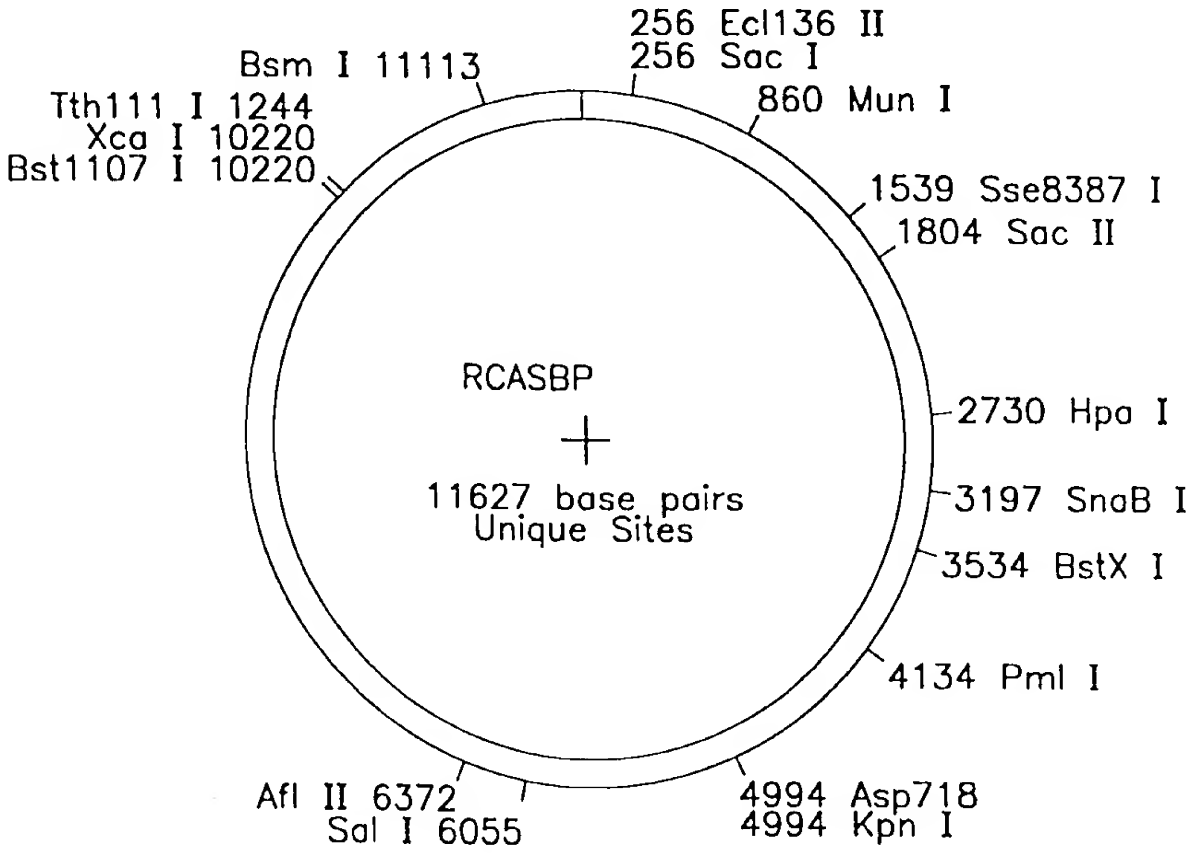


FIG. 6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256 Sac I
860 Mun I
1539 Sse8387 I
1804 Sac II
2730 Hpa I
3197 SnaB I
3534 BstX I
4134 Pml I
4994 Asp718
4994 Kpn I
Afl II 6372
Sal I 6055
Bsm I 11113
Tth111 I 1244
Xca I 10220
Bst1107 I 10220
RCASBP
11627 base pairs
Unique Sites

FIG. 7

human Yes-1 Protein amino acid sequence

"MGCIKSKENKSPAIKYRPENTPEPVSTSVSHYGAEPTTVSPCPS
SSAKGTAVNFSSLSMTPFGGSSGVTPFGGASSSFSVVPSSYPAGLTGGVTIFVALYDY
EARTTEDLSFKKGERFQIINNTEGDWWEARSIATGKNGYIPSNYVAPADSIQAEEWYF
GKMGRKDAERLLLNPGNQRGIFLVRESETTKGAYSLSIRDWDEIRGDNVKHYKIRKLD
NGGYYITTRAQFDTLQKLVKHYTEHADGLCHKLTTVCPTVKPQTQGLAKDAWEIPRES
LRLEVKLGQGCFGEVWMGTWNGTTKVAIKTLKPGTMMPEAFLQEAQIMKKLRHDKLVP
LYAVVSEEPIYIVTEFMSKGSLLDFLKEGDGKYLKLPQLVDMAAQIADGMAYIERMNY
IHRDLRAANILVGENLVCKIADFGLARLIEDNEYTARQGAKFPIKWTAPEAALYGRFT
IKSDVWSFGILQTELVTKGRVPYPGMVNREVLEQVERGYRMPCPQGCPESLHELMNLC
WKKDPDERPTFEYIQSFLEDYFTATEPQYQPGENL"

FIGURE 8

FIGURE 9

```
   1 gcggagccaa ggcacacggg tctgaccctt gggccggccc ggagcaagtg acacggaccg
  61 gtcgcctatc ctgaccacag caaagcggcc cggagcccgc ggaggggacc tgacgggggc
 121 gtaggcgccg gaaggctggg ggccccggag ccgggccggc gtggcccgag ttccggtgag
 181 cggacggcgg cgcgcgcaga tttgataatg ggctgcatta aaagtaaaga aaacaaaagt
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 301 tatggagcag aacccactac agtgtcacca tgtccgtcat cttcagcaaa gggaacagca
 361 gttaatttca gcagtctttc catgacacca tttggaggat cctcaggggt aacgcctttt
 421 ggaggtgcat cttcctcatt ttcagtggtg ccaagttcat atcctgctgg tttaacaggt
 481 ggtgttacta tatttgtggc cttatatgat tatgaagcta gaactacaga agacctttca
 541 tttaagaagg gtgaaagatt tcaaataatt aacaatacgg aaggagattg gtgggaagca
 601 agatcaatcg ctacaggaaa gaatggttat atcccgagca attatgtagc gcctgcagat
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 781 ggtgcttatt ccctttctat tcgtgattgg gatgagataa ggggtgacaa tgtgaaacac
 841 tacaaaatta ggaaacttga caatggtgga tactatatca caaccagagc acaatttgat
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 961 acaactgtgt gtccaactgt gaaacctcag actcaaggtc tagcaaaaga tgcttgggaa
1021 atccctcgag aatctttgcg actagaggtt aaactaggac aaggatgttt cggcgaagtg
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1561 aagtctgatg tctggtcatt tggaattctg caaacagaac tagtaacaaa gggccgagtg
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1741 gaccctgatg aaagaccaac atttgaatat attcagtcct tcttggaaga ctacttcact
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1861 gcacaaatct gccaaaatat aaagaacttg tgtagatttt ctacaggaat caaaagaaga
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1981 aaaccacttt tttttcccca agtattaaac tctaatgtac caatgatgaa tttatcagcg
2041 tatttcaggg tccaaacaaa atagagctaa gatactgatg acagtgtggg tgacagcatg
2101 gtaatgaagg acagtgaggc tcctgcttat ttataaatca tttcctttct ttttttcccc
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3061 gttatggctt cacattcatt gcagtgggat atggttttta tgtaaaacat ttttagaact
3121 ccagttttca aatcatgttt gaatctacat tcacttttt ttgttttctt ttttgagacg
3181 gagtctcgct ctgccgccca ggctggagtg cagtggcgcg atctcggctc actgcaagct
3241 ctgcctccca ggttcacacc attctcctgc ctcagcctcc cgagtagctg ggactacagg
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```

```
3481 atggtgg[redacted] ttttttccct tttagaatac attaaat[redacted]t tgatttgggg aggaaaactt
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3601 acatacaaaa taagactaat ttttaagagt aactcagtaa tttcaaaata cagatttgaa
3661 tagcagcatt agtggtttga gtgtctagca aaggaaaaat tgatgaataa aatgaaggtc
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3781 gcccctctat tttcaggata taattcttaa ctatcattat ttacctgatt ttaatcatca
3841 gattcgaaat tctgtgccat ggcgtatatg ttcaaattca aaccattttt aaaatgtgaa
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4081 actatgatat tgtaattttt ctaagcatat tcaaaagggt gacaaaatta cgtttatgta
4141 ctaaatctaa tcaggaaagt aaggcaggaa aagttgatgg tattcattag gttttaactg
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FIGURE 9 Con't

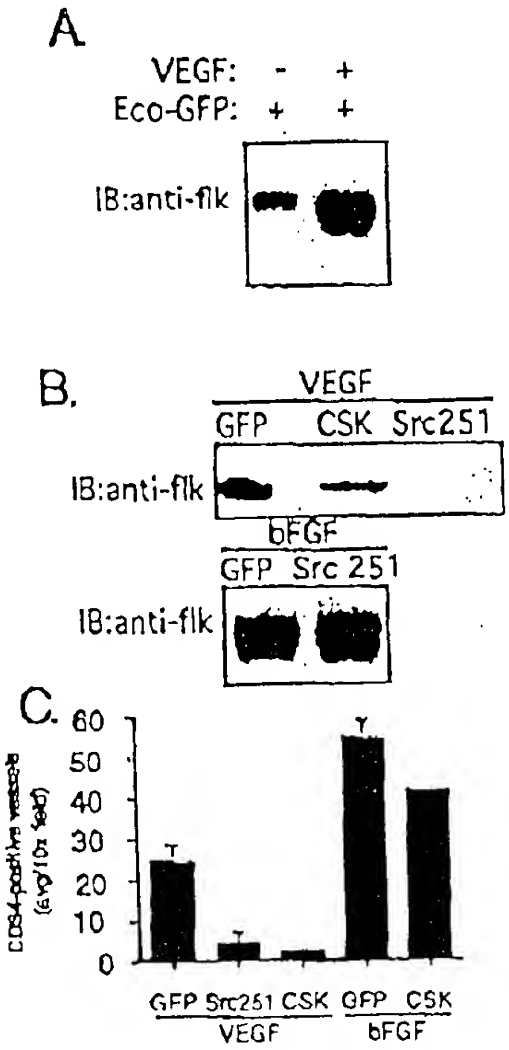

A
VEGF: - +
Eco-GFP: + +
IB:anti-flk
B.
VEGF
GFP CSK Src251
IB:anti-flk
bFGF
GFP Src 251
IB:anti-flk
C.
60
50
40
30
20
10
0
GFP Src251 CSK GFP CSK
VEGF
bFGF

FIGURE 10

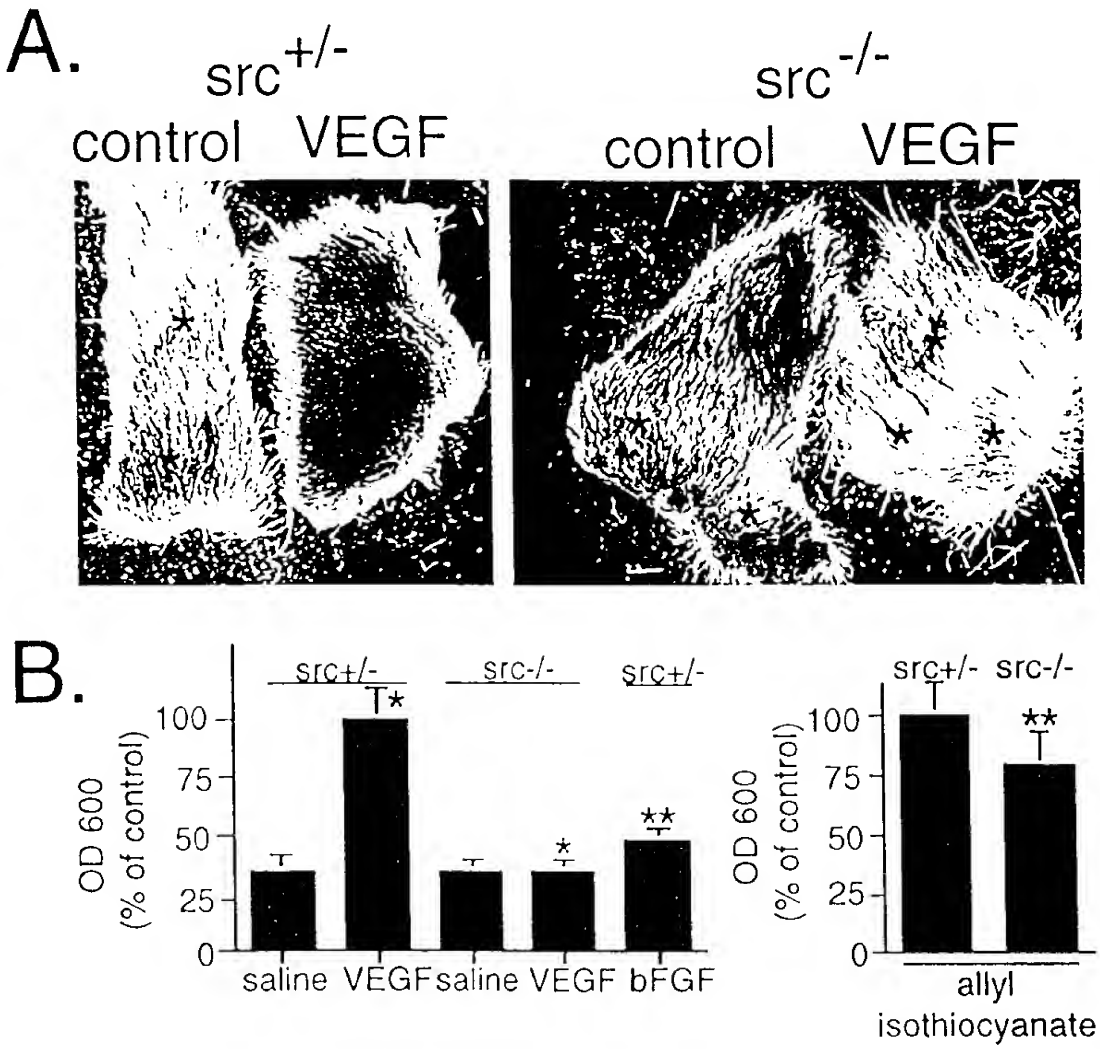
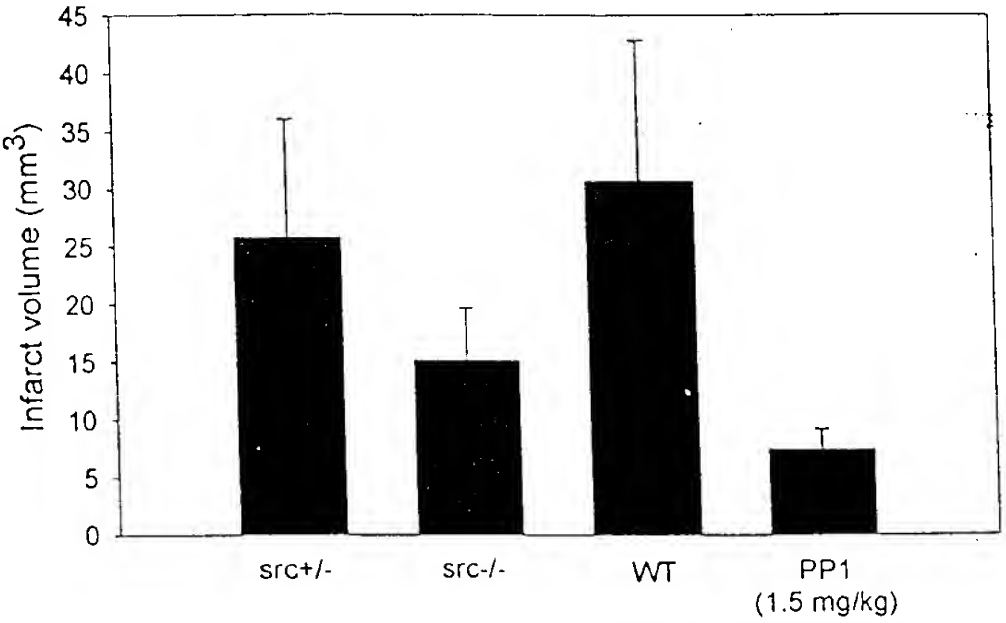

A.
src+/-
control
VEGF
src-/-
control
VEGF
B.
src+/-
src-/-
src+/-
OD 600
(% of control)
100
75
50
25
0
saline
VEGF
saline
VEGF
bFGF
src+/-
src-/-
allyl
isothiocyanate
C.
OD 600
(% of control)
100
75
50
25
0
src+/-
src-/-
fyn-/-
yes-/-

FIG. 11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40
35
30
25
20
15
10
5
0
Infarct volume (mm3)
src+/-
src-/-
WT
PP1
(1.5 mg/kg)

FIG 12

FIG 13